

WO 00/26350

PCT/US99/25820

SEQUENCE LISTING

<110> INCYTE PHARMACEUTICALS, INC.

TANG, Y. Tom
 CORLEY, Neil C.
 GUEGLER, Karl J.
 GORGONE, Gina A.
 AZIMZAI, Yalda
 KASER, Matthew R.
 YUE, Henry

<120> COENZYME A-UTILIZING ENZYMES

<130> PF-0622 PCT

<140> To Be Assigned

<141> Herewith

<150> 09/185,217; unassigned

<151> 1998-11-03; 1998-11-03

<160> 10

<170> PERL Program

<210> 1

<211> 100

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 1580751CD1

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Asn	Val	His	Glu	Lys	Lys	Leu	Gly	Asp	Lys	Val	Ala	Phe	Tyr	Trp
				20					25					30
Pro	Cys	Gln	Arg	Asp	Gln	Asp	Gly	Tyr	Tyr	Trp	Ile	Thr	Gly	Arg
				35					40					45
Ile	Asp	Asp	Met	Leu	Asn	Val	Ser	Gly	Glu	Gly	Gln	Gly	Pro	Pro
				50					55					60
Ser	His	Leu	Ile	Asn	Ser	Ala	Pro	Leu	Thr	Thr	Pro	Ser	Arg	Ser
				65					70					75
Leu	Pro	Gln	Glu	Pro	Arg	Ser	Val	Leu	Trp	Pro	Asp	His	Val	Leu
				80					85					90
Ser	Val	Ala	Phe	Ser	Ser	Gly	Pro	Arg	Phe					
				95					100					

<210> 2

<211> 159

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<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 1627889CD1

<400> 2

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          20          25          30
His Ser Glu Gln Trp Gln Leu Ser Thr Ser Gln Ile Pro Val Gln
          35          40          45
Gln Met His Leu Phe Asp Val His Asn Tyr Pro Asp Tyr Val Ser
          50          55          60
Ser Gly Gly Gly Phe Gly Pro Ala Asp Asp His Gly Tyr Gly Val
          65          70          75
Ser Tyr Ile Phe Met Gly Asp Gly Met Ile Thr Phe His Ile Ser
          80          85          90
Ser Lys Lys Ser Ser Thr Lys Thr Asp Ser His Arg Leu Gly Gln
          95          100          105
His Ile Glu Asp Ala Leu Leu Asp Val Ala Ser Leu Phe Gln Ala
          110          115          120
Gly Gln His Phe Lys Arg Arg Phe Arg Gly Ser Gly Lys Glu Asn
          125          130          135
Ser Arg His Arg Cys Gly Phe Leu Ser Arg Gln Thr Gly Ala Ser
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Lys Ala Ser Met Thr Ser Thr Asp Phe
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<210> 3

<211> 215

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<213> Homo sapiens

<220>

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<223> Incyte ID No: 1965888CD1

<400> 3

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          20          25          30
Val Arg Val Leu Leu Phe Arg Ser Gly Val Lys Gly Val Phe Cys
          35          40          45
Ala Gly Ala Asp Leu Lys Glu Arg Glu Gln Met Ser Glu Ala Glu
          50          55          60
Val Gly Val Phe Val Gln Arg Leu Arg Gly Leu Met Asn Asp Ile
          65          70          75
Ala Ser Ser Ala Val Met Gly Leu Ile Glu Thr Thr Arg Gly Leu
          80          85          90
Leu Pro Gly Ala Gly Gly Thr Gln Arg Leu Pro Arg Cys Leu Gly

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[illegible]

<210> 4

<211> 720

<212> PRT

<213> Homo sapiens

<220>

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<221> misc_feature
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<223> Incyte ID No: 2816341CD1

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His	Thr	Ile	Asn	Pro	Ile	Leu	Leu	Tyr	Phe	Ile	His	Phe	Leu	Ile
				20					25					30
Ser	Leu	Tyr	Thr	Ile	Leu	Thr	Tyr	Ile	Pro	Phe	Tyr	Phe	Phe	Ser
				35					40					45
Glu	Ser	Arg	Gln	Glu	Lys	Ser	Asn	Arg	Ile	Lys	Ala	Lys	Pro	Val
				50					55					60
Asn	Ser	Lys	Pro	Asp	Ser	Ala	Tyr	Arg	Ser	Val	Asn	Ser	Leu	Asp
				65					70					75
Gly	Leu	Ala	Ser	Val	Leu	Tyr	Pro	Gly	Cys	Asp	Thr	Leu	Asp	Lys
				80					85					90
Val	Phe	Thr	Tyr	Ala	Lys	Asn	Lys	Phe	Lys	Asn	Lys	Arg	Leu	Leu
				95					100					105
Gly	Thr	Arg	Glu	Val	Leu	Asn	Glu	Glu	Asp	Glu	Val	Gln	Pro	Asn
				110					115					120
Gly	Lys	Ile	Phe	Lys	Lys	Val	Ile	Leu	Gly	Gln	Tyr	Asn	Trp	Leu
				125					130					135
Ser	Tyr	Glu	Asp	Val	Phe	Val	Arg	Ala	Phe	Asn	Phe	Gly	Asn	Gly
				140					145					150
Leu	Gln	Met	Leu	Gly	Gln	Lys	Pro	Lys	Thr	Asn	Ile	Ala	Ile	Phe
				155					160					165
Cys	Glu	Thr	Arg	Ala	Glu	Trp	Met	Ile	Ala	Ala	Gln	Ala	Cys	Phe
				170					175					180
Met	Tyr	Asn	Phe	Gln	Leu	Val	Thr	Leu	Tyr	Ala	Thr	Leu	Gly	Gly
				185					190					195

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Pro Ala Ile Val	His Ala Leu Asn Glu	Thr Glu Val Thr Asn Ile	200	205	210
Ile Thr Ser Lys	Glu Leu Leu Gln Thr	Lys Leu Lys Asp Ile Val	215	220	225
Ser Leu Val Pro	Arg Leu Arg His Ile	Ile Thr Val Asp Gly Lys	230	235	240
Pro Pro Thr Trp	Ser Glu Phe Pro Lys	Gly Ile Ile Val His Thr	245	250	255
Met Ala Ala Val	Glu Ala Leu Gly Ala	Lys Ala Ser Met Glu Asn	260	265	270
Gln Pro His Ser	Lys Pro Leu Pro Ser	Asp Ile Ala Val Ile Met	275	280	285
Tyr Thr Ser Gly	Ser Thr Gly Leu Pro	Lys Gly Val Met Ile Ser	290	295	300
His Ser Asn Ile	Ile Ala Gly Ile Thr	Gly Met Ala Glu Arg Ile	305	310	315
Pro Glu Leu Gly	Glu Glu Asp Val Tyr	Ile Gly Tyr Leu Pro Leu	320	325	330
Ala His Val Leu	Glu Leu Ser Ala Glu	Leu Val Cys Leu Ser His	335	340	345
Gly Cys Arg Ile	Gly Tyr Ser Ser Pro	Gln Thr Leu Ala Asp Gln	350	355	360
Ser Ser Lys Ile	Lys Lys Gly Ser Lys	Gly Asp Thr Ser Met Leu	365	370	375
Lys Pro Thr Leu	Met Ala Ala Val Pro	Glu Ile Met Asp Arg Ile	380	385	390
Tyr Lys Asn Val	Met Asn Lys Val Ser	Glu Met Ser Ser Phe Gln	395	400	405
Arg Asn Leu Phe	Ile Leu Ala Tyr Asn	Tyr Lys Met Glu Gln Ile	410	415	420
Ser Lys Gly Arg	Asn Thr Pro Leu Cys	Asp Ser Phe Val Phe Arg	425	430	435
Lys Val Arg Ser	Leu Leu Gly Gly Asn	Ile Arg Leu Leu Leu Cys	440	445	450
Gly Gly Ala Pro	Leu Ser Ala Thr Thr	Gln Arg Phe Met Asn Ile	455	460	465
Cys Phe Cys Cys	Pro Val Gly Gln Gly	Tyr Gly Leu Thr Glu Ser	470	475	480
Ala Gly Ala Gly	Thr Ile Ser Glu Val	Trp Asp Tyr Asn Thr Gly	485	490	495
Arg Val Gly Ala	Pro Leu Val Cys Cys	Glu Ile Lys Leu Lys Asn	500	505	510
Trp Glu Glu Gly	Gly Tyr Phe Asn Thr	Asp Lys Pro His Pro Arg	515	520	525
Gly Glu Ile Leu	Ile Gly Gly Gln Ser	Val Thr Met Gly Tyr Tyr	530	535	540
Lys Asn Glu Ala	Lys Thr Lys Ala Asp	Phe Phe Glu Asp Glu Asn	545	550	555
Gly Gln Arg Trp	Leu Cys Thr Gly Asp	Ile Gly Glu Phe Glu Pro	560	565	570
Asp Gly Cys Leu	Lys Ile Ile Asp Arg	Lys Lys Asp Leu Val Lys	575	580	585
Leu Gln Ala Gly	Glu Tyr Val Ser Leu	Gly Lys Val Glu Ala Ala	590	595	600
Leu Lys Asn Leu	Pro Leu Val Asp Asn	Ile Cys Ala Tyr Ala Asn			

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	605		610		615
Ser Tyr His Ser	Tyr Val Ile Gly Phe	Val Val Pro Asn Gln	Lys		
	620		625		630
Glu Leu Thr Glu	Leu Ala Arg Lys Lys	Gly Leu Lys Gly Thr	Trp		
	635		640		645
Glu Glu Leu Cys	Asn Ser Cys Glu Met	Glu Asn Glu Leu Leu	Lys		
	650		655		660
Val Leu Ser Glu	Ala Ala Ile Ser Ala	Ser Leu Glu Lys Phe	Glu		
	665		670		675
Ile Leu Val Lys	Ile Arg Leu Ser Pro	Glu Pro Trp Thr Pro	Glu		
	680		685		690
Thr Gly Leu Val	Thr Asp Ala Phe Lys	Leu Lys Arg Lys Glu	Leu		
	695		700		705
Lys Thr His Tyr	Gln Ala Asp Ile Glu	Arg Met Tyr Gly Arg	Lys		
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<210> 5

<211> 456

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3324214CD1

<400> 5

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Gly Ile Ser Leu Thr	Val Leu Phe Thr Leu	Leu Leu Val Phe Ile	
	20	25	30
Ile Val Pro Ala Ile	Phe Gly Val Ser Phe	Gly Ile Arg Lys Leu	
	35	40	45
Tyr Met Lys Ser Leu	Leu Lys Ile Phe Ala	Trp Ala Thr Leu Arg	
	50	55	60
Met Glu Arg Gly Ala	Lys Glu Lys Asn His	Gln Leu Tyr Lys Pro	
	65	70	75
Tyr Thr Asn Gly Ile	Ile Ala Lys Asp Pro	Thr Ser Leu Glu Glu	
	80	85	90
Glu Ile Lys Glu Ile	Arg Arg Ser Gly Ser	Ser Lys Ala Leu Asp	
	95	100	105
Asn Thr Pro Glu Phe	Glu Leu Ser Asp Ile	Phe Tyr Phe Cys Arg	
	110	115	120
Lys Gly Met Glu Thr	Ile Met Asp Asp Glu	Val Thr Lys Arg Phe	
	125	130	135
Ser Ala Glu Glu Leu	Glu Ser Trp Asn Leu	Leu Ser Arg Thr Asn	
	140	145	150
Tyr Asn Phe Gln Tyr	Ile Ser Leu Arg Leu	Thr Val Leu Trp Gly	
	155	160	165
Leu Gly Val Leu Ile	Arg Tyr Cys Phe Leu	Leu Pro Leu Arg Ile	
	170	175	180
Ala Leu Ala Phe Thr	Gly Ile Ser Leu Leu	Val Val Gly Thr Thr	
	185	190	195
Val Val Gly Tyr Leu	Pro Asn Gly Arg Phe	Lys Glu Phe Met Ser	
	200	205	210

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Lys His Val His	Leu Met Cys Tyr Arg	Ile Cys Val Arg Ala	Leu
215		220	225
Thr Ala Ile Ile	Thr Tyr His Asp Arg	Glu Asn Arg Pro Arg	Asn
230		235	240
Gly Gly Ile Cys	Val Ala Asn His Thr	Ser Pro Ile Asp Val	Ile
245		250	255
Ile Leu Ala Ser	Asp Gly Tyr Tyr Ala	Met Val Gly Gln Val	His
260		265	270
Gly Gly Leu Met	Gly Val Ile Gln Arg	Ala Met Val Lys Ala	Cys
275		280	285
Pro His Val Trp	Phe Glu Arg Ser Glu	Val Lys Asp Arg His	Leu
290		295	300
Val Ala Lys Arg	Leu Thr Glu His Val	Gln Asp Lys Ser Lys	Leu
305		310	315
Pro Ile Leu Ile	Phe Pro Glu Gly Thr	Cys Ile Asn Asn Thr	Ser
320		325	330
Val Met Met Phe	Lys Lys Gly Ser Phe	Glu Ile Gly Ala Thr	Val
335		340	345
Tyr Pro Val Ala	Ile Lys Tyr Asp Pro	Gln Phe Gly Asp Ala	Phe
350		355	360
Trp Asn Ser Ser	Lys Tyr Gly Met Val	Thr Tyr Leu Leu Arg	Met
365		370	375
Met Thr Ser Trp	Ala Ile Val Cys Ser	Val Trp Tyr Leu Pro	Pro
380		385	390
Met Thr Arg Glu	Ala Asp Glu Asp Ala	Val Gln Phe Ala Asn	Arg
395		400	405
Val Lys Ser Ala	Ile Ala Arg Gln Gly	Gly Leu Val Asp Leu	Leu
410		415	420
Trp Asp Gly Gly	Leu Lys Arg Glu Lys	Val Lys Asp Thr Phe	Lys
425		430	435
Glu Glu Gln Gln	Lys Leu Tyr Ser Lys	Met Ile Val Gly Asn	His
440		445	450
Lys Asp Arg Ser	Arg Ser		
455			

<210> 6

<211> 687

<212> DNA

<213> Homo sapiens

<220>

<221> unsure

<222> 63

<223> a or g or c or t, unknown, or other

<220>

<221> misc_feature

<223> Incyte ID No: 1580751CB1

<400> 6

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cgaggtcagc cgctccgcgc acgtcccctc gctgcagcgc taccgcgagc tgcaccggcg 180
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tgagaaaaag cttggagata aagttgcttt ttactggcct tgccagcggg accaggatgg 420
ctattactgg atcaactggca ggattgatga catgctcaat gtatctggtg agggccaggg 480
gccaccttcc catcttatta actctgctcc tctgacaaca cccagccgaa gccttccgca 540
agagcccagg agtgtccttt ggccagacca tgtactaagt gtagcattca gttctggggc 600
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<210> 7

<211> 1803

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 1627889CB1

<400> 7

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gtgctcagac taaagctttt ctttatacgg gggttcatgt tggtaaaaat ccattgaatt 180
atatacgtag gatttttgaa cttatttcta taaagtttta tatttcaata aaaagcttaa 240
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<210> 8

<211> 1340

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<212> DNA

<213> Homo sapiens

<220>

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<223> Incyte ID No: 1965888CB1

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gcgggtcccg accaagttta caagtccttc cactttctct ctgaggcaga aagagcaagg 180
gtttttctct ccattttatg gttgggaaaa ttgaggcctg cctgagtgtg tgacttgttg 240
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<210> 9

<211> 4027

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2816341CB1

<400> 9

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gggcggcggc gcggggcgtg aacgctctgg ggctcagcca ggctgcgcg ggcccgatgc 180
cggaggaacc cggactccgg cgtagcggtt ttgacacaag ggcgcataat tcaaaagcac 240
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cagctgtaac atttggcacc ctcaagaagc gctggctcctg tgtcacacca ccttagcctc 360
ttgatcgagg aagattctcg ctgaagtctg ttaattctac tttttgagta cttatgaata 420
accacgtgtc ttcaaaacca tctaccatga agctaaaaca taccatcaac cctattcttt 480
tatattttat acattttcta atatcacttt atactatttt aacatacatt ccgttttatt 540
ttttctccga gtcaagacaa gaaaaatcaa accgaattaa agcaaagcct gttaaattcaa 600
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tttttaaaaa	ggttattctt	ggacagtata	attggctttc	ctatgaagat	gtctttgttc	840
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